HU: MGSDVRDLNALLPAVPSLGGGGGCALPVSGAAQWAPVLDFAPPGASAYGSL MO: MGSDVRDLNALLPAVSSLGGGGGCGLPVSGAAQWAPVLDFAPPGASAYGSL

HU:GGPAPPPAPPPPPPPPPPHSFIKQEPSWGGAEPHEEQCLSAFTVHFSGQFTGTAG MO:GGPAPPPAPPPPPPPPPPHSFIKQEPSWGGAEPHEEQCLSAFTLHFSGQFTGTAG

HU: ACRYGPFGPPPPSQASSGQARMFPNAPYLPSCLESQPAIRNQGYSTVTFDGTPS MO: ACRYGPFGPPPPSQASSGQARMFPNAPYLPSCLESQPTIRNQGYSTVTFDGAPS

HU: YGHTPSHHAAQFPNHSFKHEDPMGQQGSLGEQQYSVPPPVYGCHTPTDSCTG MO: YGHTPSHHAAQFPNHSFKHEDPMGQQGSLGEQQYSVPPPVYGCHTPTDSCTG

HU: SQALLLRTPYSSDNLYQMTSQLECMTWNQMNLGATLKGVAAGSSSSVKWTE MO: SQALLLRTPYSSDNLYQMTSQLECMTWNQMNLGATLKGMAAGSSSSVKWTE

HU:GQSNHSTGYESDNHTTPILCGAQYRIHTHGVFRGIQDVRRVPGVAPTLVRSAS MO:GQSNHGIGYESDNHTAPILCGAQYRIHTHGVFRGIQDVRRVSGVAPTLVRSAS

HU: ETSEKRPFMCAYPGCNKRYFKLSHLQMHSRKHTGEKPYQCDFKDCERRFSR MO: ETSEKRPFMCAYPGCNKRYFKLSHLQMHSRKHTGEKPYQCDFKDCERRFSR

HU: SDQLKRHQRRHTGVKPFQCKTCQRKFSRSDHLKTHTRTHTGKTSEKPFSCR MO: SDQLKRHQRRHTGVKPFQCKTCQRKFSRSDHLKTHTRTHTGKTSEKPFSCR

HU:WPSCQKKFARSDELVRHHNMHQRNMTKLQLAL MO:WHSCQKKFARSDELVRHHNMHQRNMTKLHVAL



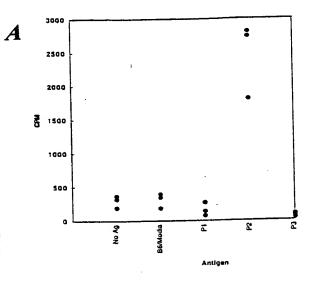
FIG. 2

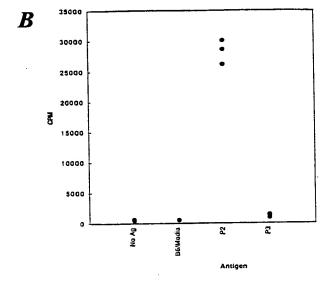


FIG. 3



FIG. 4





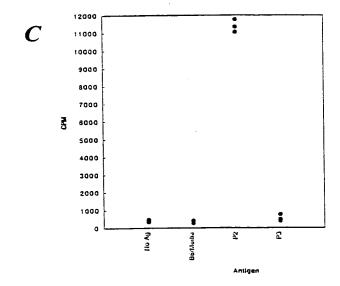
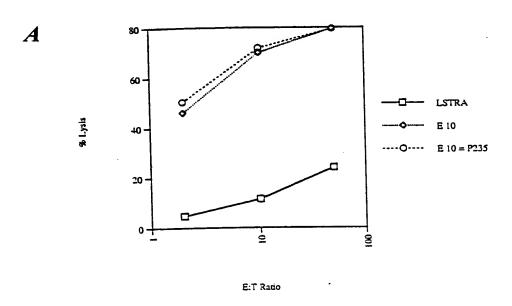


FIG. 5A-5C



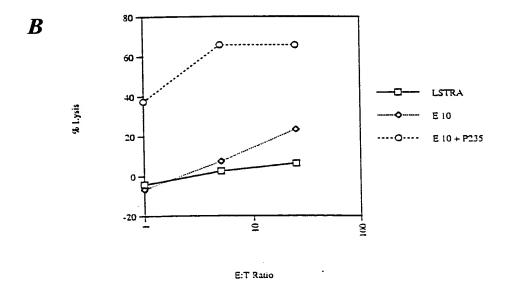
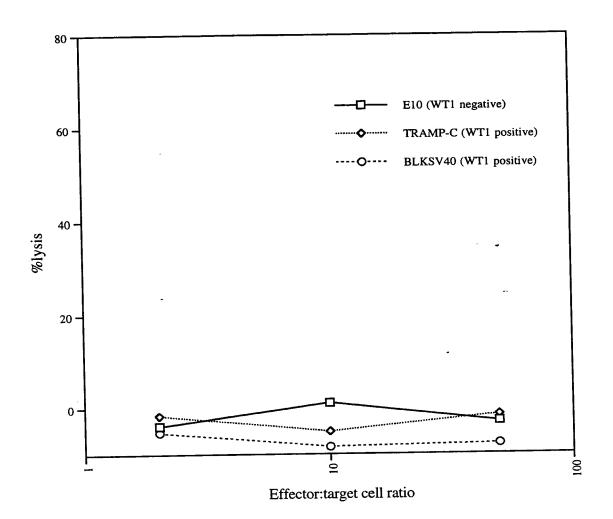


FIG. 6A and 6B



. FIG. 7A

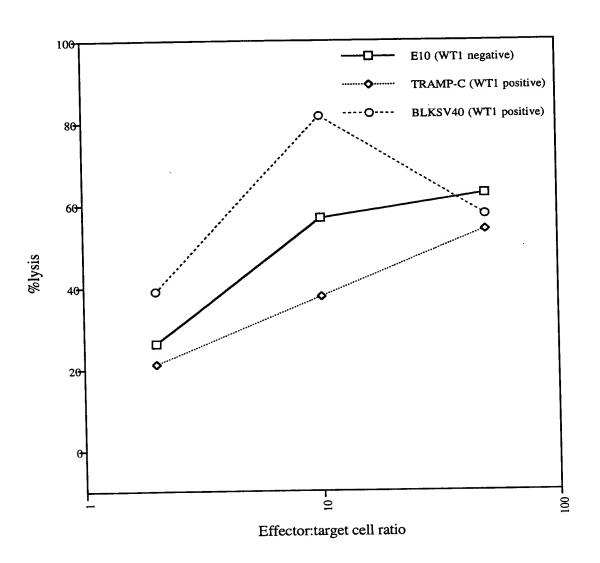


FIG. 7B

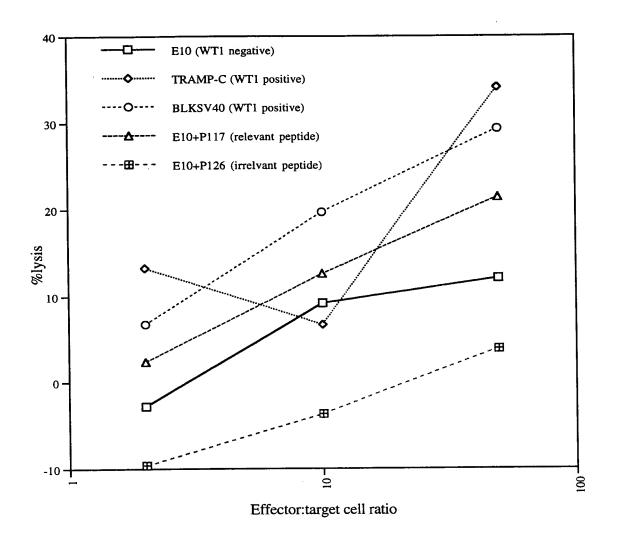


FIG. 7C

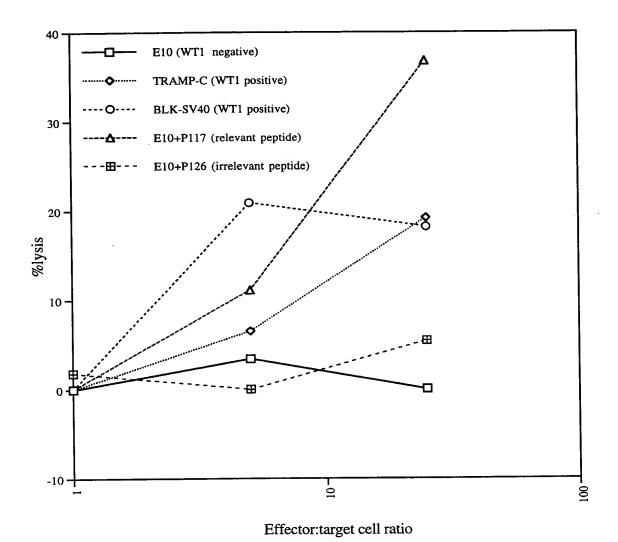


FIG. 7D

5 MGSDVRD	AAAAA	AAAA	AAAAA	GCALP'	. AAA.	QWAPV AAA	 	GASA AAA				 		
														
80 PSWGGAI		AA	TVHFS A	GQFTG AAA R	TAGAC A RRR	RYGPF		SQASS · · · · · · · · · ·	RRI	FPNAP	YLPSC:	LESQP AA	AIRNQ	
	 								• • • • •		• • • • •		• • • • •	
155 TVTFDG	TPSYGH	TPSHH	AAAA.		HEDPM	IGQQGS		YSVPF	PVYGC A	HTPTC AAAAA	SCTGS	QALLI DDDDI	RTPYS	
230 LYQMTS AAAAAA	235 QLECMI AA	240 WNQMI	245 NLGATI A	250 LKGVAA AA. AAA RRRRRE	255 AGSSSS A	260 SVKWTE	265 EGQSNH	270 ISTGYE	275 ESDNHT	280 TPILC	285 CGAQYR	290 EIHTHO AA	295 GVFRGI AAAAA RRRR	300 QDV AAA
305 RRVPGV		215	220	225	330	335	340	345	350	355	360	365	370	375
AAAAA.	APTLVE AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	AAAA. 	AA		 	 .RRRR		· · · · · ·		 				
	385 KPFQCK'	AAA	FSRSD A.AAA	AAA 	TRTHT	AAAA.	KPFSCI	A	QKKFAI AA RRRI	RSDEL .AAAA RRR	RR			

P	10 DLNALI AAAAA	LPAVSS AAAAA	LGGGG	4	VSGAA AAA	AQWAPV AAAA	LDFAF	PGASA AAA	YGSLG	GPAPI AAAA	PAPPE			
				100	105	110	115	120	125	130	135	140	145	150
	YEPHEE			AAA	RRR.			 וממממכ	AAA RF ODDD	RRRR.		 		
155 TVTFD	160 GAPSYG	165 HTPSHI	170 HAAQF AAAAA	175 PNHSFF	180 KHEDP	185 MGQQGS	190 SLGEQO	195 QYSVP1	200 PPVYG0	205 CHTPT AAAAA	210 DSCTG: A. ~	215 SQALLI	220 LRTPYS	225 SDN .AA
AAAAA	SQLECM AAA	TWNQM	NLGAT A RRR	AA AA RRRRRI	AGSSS A RR	SVKWT	EGQSN 	HGIGY	ESDNH	TAPIL		RIHTHO	RRRR	QDV AAA · · ·
305 RRVSG AAAAA	310 VAPTLV AAAA	315 /RSASE AAAAA	320 TSEKF	325 RPFMCA	330 YPGCN	335 IKRYFK 	340 LSHLQ	345 MHSRK	350 HTGEK	355 PYQCE	360 DFKDCE	365 ERRFSR AAA.A	370 SDQLKF AAAAAA	375 RHQR AAA.
380 RHTGV		390 KTCQRF	395 KFSRSI AA.AA	400 OHLKTH	405 TRTH:	410 FGKTSE .AAAA.	415 KPFSC	420 CRWHSC	425 CQKKFA .AA	430 RSDEI .AAAZ	435 LVRHHN AAAAA RRR	440 MHQRN AAA	445 MTKLHV	450 VAL